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TECH CENTER 1600/2900

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1600

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/721,904B

DATE: 03/20/2003 P.6
TIME: 16:03:44

Input Set : A:\seqlist.asc.txt

Output Set: N:\CRF4\03202003\I721904B.raw

```
3 <110> APPLICANT: JULIUS, Michael H.
4     FILIPP, Dominik
6 <120> TITLE OF INVENTION: THE INDUCTION OF ANTIBIOTIC PROTEINS AND PEPTIDES BY
7     LAIT/sCD14-PROTEIN
9 <130> FILE REFERENCE: 47841/00063
11 <140> CURRENT APPLICATION NUMBER: US 09/721,904B
12 <141> CURRENT FILING DATE: 2000-11-27
14 <150> PRIOR APPLICATION NUMBER: PCT/CA99/00482
15 <151> PRIOR FILING DATE: 1999-05-27
17 <150> PRIOR APPLICATION NUMBER: US 60/086,884
18 <151> PRIOR FILING DATE: 1998-05-27
20 <160> NUMBER OF SEQ ID NOS: 11
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25 <211> LENGTH: 1122
26 <212> TYPE: DNA
27 <213> ORGANISM: bovine
29 <400> SEQUENCE: 1
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32 gacacaacag aaccctgcga gctggacgac gacgatttcc gttgtgtctg caacttcacg      120
34 gatccgaagc ctgactggtc tagcgccgtt cagtgtatgg ttgccgtcga ggtggagatc      180
36 agtgccggcg gccgcagcct ggaacagttt ctcaaggagg ccgacaccaa cccgaagcag      240
38 tatgctgaca caatcaaggc tctgcgcgtt cggcgactca agctgggcgc tgcacagggt      300
40 cctgctcagc ttctggtcgc cgttctgcgc gcgctcggtt actctcgtct caaggaactg      360
42 acgcttgagg acctggaggt aaccggccca acgccccga cgctctgga agccgctggg      420
44 cctgcgctca ccacctcag tctgcgtaac gtatcgtgga caacaggagg tgcttggtc      480
46 ggcgaactgc agcagtggct caagcctggg ctcaagggtg tgaacattgc ccaagcacac      540
48 tcgcttgctt ttccgtgcgc agggctctcc accttcgagg cgctcaccac cctagacctg      600
50 tctgacaatc ccagtctcgg cgacacgggg ctgatggcag ctctctgtcc gaacaagttc      660
52 cgggccctcc aatatctagc gctacgcaac gcggggatgg agacgccgag cggcgtgtgc      720
54 gcggcgctgg cggcagcgag ggtgcagccc caaagcctgg acctcagcca caactcgctg      780
56 cgcgtcaccg ccccggtgct taccgatgt gtctggccca gtgcactaag gtctctcaat      840
58 ttgtcgcttc ctgggctgga gcaagtgcct aagggaactgc cccctaagct cagcgtgctt      900
60 gatctcagct gcaacaagct aagcagggag ccgcggcgag acgagctgcc cgaggtaaat      960
62 gacctgactc tggacggaaa tccctttctg gacctggag ccctccagca ccaaaatgac      1020
64 ccgatgatct ccggcgtggt ccagcctgt gcgcgttctg ccttgaccat gggggtgtca      1080
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70 <211> LENGTH: 1128
71 <212> TYPE: DNA
72 <213> ORGANISM: human
74 <400> SEQUENCE: 2
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77 acgccagaac cttgtgagct ggacgatgaa gatttccgct gcgtctgcaa cttctccgaa      120
79 cctcagcccg actggtccga agccttccag tgtgtgtctg cagtagaggt ggagatccat      180
81 gccggcggtc tcaacctaga gccgtttcta aagcgcgctg atgcggacgc cgacccgcgg      240
83 cagtatgctg acacggtcaa ggctctccgc gtgcggcggc tcacagtggg agccgcacag      300
85 gttcctgctc agctactggt aggcgccctg cgtgtgctag cgtactcccg cctcaaggaa      360
87 ctgacgctcg aggacctaaa gataaccggc accatgcctc cgctgcctct ggaagccaca      420
89 ggacttgcac tttccagctt gcgcctacgc aacgtgtcgt gggcgacagg gcgttcttgg      480
91 ctgcgccgagc tgcagcagtg gctcaagcca ggcctcaagg tactgagcat tgcccaagca      540
93 cactgcgctg ccttttcctg cgaacagggt cgcgcccttc cggcccttac cagcctagac      600
95 ctgtctgaca atcctggact gggcgaacgc ggactgatgg cggtctctctg tccccacaag      660
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99 tgcgccgcac tggcggcggc aggtgtgcag cccacagacc tagacctcag ccacaactcg      780
101 ctgcgcgcca ccgtaaacc tagcgctccg agatgcatgt ggtccagcgc cctgaactcc      840
103 ctcaatctgt cgctcgctgg gctggaacag gtgcctaaag gactgccagc caagctcaga      900
105 gtgctcgatc tcagctgcaa cagactgaac agggcgccgc agcctgacga gctgcccgag      960
107 gtggataacc tgacactgga cgggaatccc ttctggtcc ctggaactgc cctccccac      1020
109 gagggtcaca tgaactccgg cgtggtccca gcctgtgcac gttcgaccct gtcggtgggg      1080
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115 <211> LENGTH: 1101
116 <212> TYPE: DNA
117 <213> ORGANISM: murine
119 <400> SEQUENCE: 3
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124 tgggtccagcg ctttcaattg tttgggggcg gcagatgtgg aattgtacgg cggcgcccg      180
126 agcctggaat accttctaaa gcgtgtggac acggaagcag atctggggca gttcactgat      240
128 attatcaagt ctctgtcctt aaagcggctt acggtgcggg ccgcgcggat tctagtctgg      300
130 attctattcg gagccctgcg tgtgtctggg atttccggcc tccaggaaact gactcttgaa      360
132 aatctcgagg taaccggcac cgcgccgcca cgcttctgg aagccaccgg acccgatctc      420
134 aacatcttga acctccgcaa cgtgtcgtgg gcaacaagg atgcctggct cgcagaactg      480
136 cagcagtggc taaagcctgg actcaaggta ctgagtattg cccaagcaca ctactcaac      540
138 ttttcctgcg aacaggctcg cgtcttccct gccctctcca ccttagacct gtctgacaat      600
140 cctgaattgg gcgagagagg actgatctca gccctctgtc ccctcaagtt cccgaccctc      660
142 caagtttttag cgctgcgtaa cgcggggatg gagacgcca gcggcgtgtg ctctgcgctg      720
144 gccgcagcaa gggtagagct gcaaggacta gacctagtc acaattcact gcgggatgct      780
146 gcaggcgctc cgagttgtga ctggcccagt cagctaaact cgctcaatct gtctttcact      840
148 gggctgaagc aggtacctaa agggctgcc aaccaagctca gcgtgctgga tctcagttac      900
150 aacaggctgg ataggaaccc tagcccagat gagctgcccc aagtggggaa cctgtcactt      960
152 aaaggaaatc cttttttgga ctctgaatcc cactcgga agtttaactc tggcgtagtc      1020
154 accgccggag ctccatcatc ccaagcagtg gcctgtcag gaactctggc tttgtccta      1080
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159 <210> SEQ ID NO: 4
160 <211> LENGTH: 373
161 <212> TYPE: PRT
162 <213> ORGANISM: bovine
164 <400> SEQUENCE: 4
165 Met Val Cys Val Pro Tyr Leu Leu Leu Leu Leu Leu Pro Ser Leu Leu
166 1 5 10 15

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168 Arg Val Ser Ala Asp Thr Thr Glu Pro Cys Glu Leu Asp Asp Asp Asp
169          20          25          30
171 Phe Arg Cys Val Cys Asn Phe Thr Asp Pro Lys Pro Asp Trp Ser Ser
172          35          40          45
174 Ala Val Gln Cys Met Val Ala Val Glu Val Glu Ile Ser Ala Gly Gly
175          50          55          60
177 Arg Ser Leu Glu Gln Phe Leu Lys Gly Ala Asp Thr Asn Pro Lys Gln
178 65          70          75          80
180 Tyr Ala Asp Thr Ile Lys Ala Leu Arg Val Arg Arg Leu Lys Leu Gly
181          85          90          95
183 Ala Ala Gln Val Pro Ala Gln Leu Leu Val Ala Val Leu Arg Ala Leu
184          100          105          110
186 Gly Tyr Ser Arg Leu Lys Glu Leu Thr Leu Glu Asp Leu Glu Val Thr
187          115          120          125
189 Gly Pro Thr Pro Pro Thr Pro Leu Glu Ala Ala Gly Pro Ala Leu Thr
190          130          135          140
192 Thr Leu Ser Leu Arg Asn Val Ser Trp Thr Thr Gly Gly Ala Trp Leu
193 145          150          155          160
195 Gly Glu Leu Gln Gln Trp Leu Lys Pro Gly Leu Arg Val Leu Asn Ile
196          165          170          175
198 Ala Gln Ala His Ser Leu Ala Phe Pro Cys Ala Gly Leu Ser Thr Phe
199          180          185          190
201 Glu Ala Leu Thr Thr Leu Asp Leu Ser Asp Asn Pro Ser Leu Gly Asp
202          195          200          205
204 Thr Gly Leu Met Ala Ala Leu Cys Pro Asn Lys Phe Pro Ala Leu Gln
205          210          215          220
207 Tyr Leu Ala Leu Arg Asn Ala Gly Met Glu Thr Pro Ser Gly Val Cys
208 225          230          235          240
210 Ala Ala Leu Ala Ala Ala Arg Val Gln Pro Gln Ser Leu Asp Leu Ser
211          245          250          255
213 His Asn Ser Leu Arg Val Thr Ala Pro Gly Ala Thr Arg Cys Val Trp
214          260          265          270
216 Pro Ser Ala Leu Arg Ser Leu Asn Leu Ser Phe Ala Gly Leu Glu Gln
217          275          280          285
219 Val Pro Lys Gly Leu Pro Pro Lys Leu Ser Val Leu Asp Leu Ser Cys
220          290          295          300
222 Asn Lys Leu Ser Arg Glu Pro Arg Arg Asp Glu Leu Pro Glu Val Asn
223 305          310          315          320
225 Asp Leu Thr Leu Asp Gly Asn Pro Phe Leu Asp Pro Gly Ala Leu Gln
226          325          330          335
228 His Gln Asn Asp Pro Met Ile Ser Gly Val Val Pro Ala Cys Ala Arg
229          340          345          350
231 Ser Ala Leu Thr Met Gly Val Ser Gly Ala Leu Ala Leu Leu Gln Gly
232          355          360          365
234 Ala Arg Gly Phe Ala
235          370
238 <210> SEQ ID NO: 5
239 <211> LENGTH: 375
240 <212> TYPE: PRT

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RAW SEQUENCE LISTING

DATE: 03/20/2003

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Output Set: N:\CRF4\03202003\I721904B.raw

241 <213> ORGANISM: human

243 <400> SEQUENCE: 5

244 Met Glu Arg Ala Ser Cys Leu Leu Leu Leu Leu Leu Pro Leu Val His

245 1 5 10 15

247 Val Ser Ala Thr Thr Pro Glu Pro Cys Glu Leu Asp Asp Glu Asp Phe

248 20 25 30

250 Arg Cys Val Cys Asn Phe Ser Glu Pro Gln Pro Asp Trp Ser Glu Ala

251 35 40 45

253 Phe Gln Cys Val Ser Ala Val Glu Val Glu Ile His Ala Gly Gly Leu

254 50 55 60

256 Asn Leu Glu Pro Phe Leu Lys Arg Val Asp Ala Asp Ala Asp Pro Arg

257 65 70 75 80

259 Gln Tyr Ala Asp Thr Val Lys Ala Leu Arg Val Arg Arg Leu Thr Val

260 85 90 95

262 Gly Ala Ala Gln Val Pro Ala Gln Leu Leu Val Gly Ala Leu Arg Val

263 100 105 110

265 Leu Ala Tyr Ser Arg Leu Lys Glu Leu Thr Leu Glu Asp Leu Lys Ile

266 115 120 125

268 Thr Gly Thr Met Pro Pro Leu Pro Leu Glu Ala Thr Gly Leu Ala Leu

269 130 135 140

271 Ser Ser Leu Arg Leu Arg Asn Val Ser Trp Ala Thr Gly Arg Ser Trp

272 145 150 155 160

274 Leu Ala Glu Leu Gln Gln Trp Leu Lys Pro Gly Leu Lys Val Leu Ser

275 165 170 175

277 Ile Ala Gln Ala His Ser Pro Ala Phe Ser Tyr Glu Gln Val Arg Ala

278 180 185 190

280 Phe Pro Ala Leu Thr Ser Leu Asp Leu Ser Asp Asn Pro Gly Leu Gly

281 195 200 205

283 Glu Arg Gly Leu Met Ala Ala Leu Cys Pro His Lys Phe Pro Ala Ile

284 210 215 220

286 Gln Asn Leu Ala Leu Arg Asn Thr Gly Met Glu Thr Pro Thr Gly Val

287 225 230 235 240

289 Cys Ala Ala Leu Ala Ala Ala Gly Val Gln Pro His Ser Leu Asp Leu

290 245 250 255

292 Ser His Asn Ser Leu Arg Ala Thr Val Asn Pro Ser Ala Pro Arg Cys

293 260 265 270

295 Met Trp Ser Ser Ala Leu Asn Ser Leu Asn Leu Ser Phe Ala Gly Leu

296 275 280 285

298 Glu Gln Val Pro Lys Gly Leu Pro Ala Lys Leu Arg Val Leu Asp Leu

299 290 295 300

301 Ser Cys Asn Arg Leu Asn Arg Ala Pro Gln Pro Asp Glu Leu Pro Glu

302 305 310 315 320

304 Val Asp Asn Leu Thr Leu Asp Gly Asn Pro Phe Leu Val Pro Gly Thr

305 325 330 335

307 Ala Leu Pro His Glu Gly Ser Met Asn Ser Gly Val Val Pro Ala Cys

308 340 345 350

310 Ala Arg Ser Thr Leu Ser Val Gly Val Ser Gly Thr Leu Val Leu Leu

311 355 360 365

313 Gln Gly Ala Arg Gly Phe Ala

RAW SEQUENCE LISTING

DATE: 03/20/2003

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TIME: 16:03:44

Input Set : A:\seqlist.asc.txt

Output Set: N:\CRF4\03202003\I721904B.raw

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314      370      375
317 <210> SEQ ID NO: 6
318 <211> LENGTH: 366
319 <212> TYPE: PRT
320 <213> ORGANISM: murine
322 <400> SEQUENCE: 6
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324 1      5      10      15
326 Pro Ala Pro Pro Glu Pro Cys Glu Leu Asp Glu Glu Ser Cys Ser Cys
327      20      25      30
329 Asn Phe Ser Asp Pro Lys Pro Asp Trp Ser Ser Ala Phe Asn Cys Leu
330      35      40      45
332 Gly Ala Ala Asp Val Glu Leu Tyr Gly Gly Gly Arg Ser Leu Glu Tyr
333      50      55      60
335 Leu Leu Lys Arg Val Asp Thr Glu Ala Asp Leu Gly Gln Phe Thr Asp
336 65      70      75      80
338 Ile Ile Lys Ser Leu Ser Leu Lys Arg Leu Thr Val Arg Ala Ala Arg
339      85      90      95
341 Ile Pro Ser Arg Ile Leu Phe Gly Ala Leu Arg Val Leu Gly Ile Ser
342      100     105     110
344 Gly Leu Gln Glu Leu Thr Leu Glu Asn Leu Glu Val Thr Gly Thr Ala
345      115     120     125
347 Pro Pro Pro Leu Leu Glu Ala Thr Gly Pro Asp Leu Asn Ile Leu Asn
348      130     135     140
350 Leu Arg Asn Val Ser Trp Ala Thr Arg Asp Ala Trp Leu Ala Glu Leu
351 145     150     155     160
353 Gln Gln Trp Leu Lys Pro Gly Leu Lys Val Leu Ser Ile Ala Gln Ala
354      165     170     175
356 His Ser Leu Asn Phe Ser Cys Glu Gln Val Arg Val Phe Pro Ala Leu
357      180     185     190
359 Ser Thr Leu Asp Leu Ser Asp Asn Pro Glu Leu Gly Glu Arg Gly Leu
360      195     200     205
362 Ile Ser Ala Leu Cys Pro Leu Lys Phe Pro Thr Leu Gln Val Leu Ala
363      210     215     220
365 Leu Arg Asn Ala Gly Met Glu Thr Pro Ser Gly Val Cys Ser Ala Leu
366 225     230     235     240
368 Ala Ala Ala Arg Val Gln Leu Gln Gly Leu Asp Leu Ser His Asn Ser
369      245     250     255
371 Leu Arg Asp Ala Ala Gly Ala Pro Ser Cys Asp Trp Pro Ser Gln Leu
372      260     265     270
374 Asn Ser Leu Asn Leu Ser Phe Thr Gly Leu Lys Gln Val Pro Lys Gly
375      275     280     285
377 Leu Pro Ala Lys Leu Ser Val Leu Asp Leu Ser Tyr Asn Arg Leu Asp
378      290     295     300
380 Arg Asn Pro Ser Pro Asp Glu Leu Pro Gln Val Gly Asn Leu Ser Leu
381 305     310     315     320
383 Lys Gly Asn Pro Phe Leu Asp Ser Glu Ser His Ser Glu Lys Phe Asn
384      325     330     335
386 Ser Gly Val Val Thr Ala Gly Ala Pro Ser Ser Gln Ala Val Ala Leu

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RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/09/721,904B

DATE: 03/20/2003
TIME: 16:03:45

Input Set : A:\seqlist.asc.txt
Output Set: N:\CRF4\03202003\I721904B.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:7; Xaa Pos. 14,265,266,267,269

VERIFICATION SUMMARY

PATENT APPLICATION: **US/09/721,904B**

DATE: 03/20/2003

TIME: 16:03:45

Input Set : **A:\seqlist.asc.txt**

Output Set: **N:\CRF4\03202003\I721904B.raw**

L:120 M:112 C: (48) String data converted to lower case,
M:112 Repeated in SeqNo=3
L:414 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:0
M:341 Repeated in SeqNo=7